SEQUENCE LISTING

<110> The Scripps Research Institute
Deiters, Alexander
Cropp, T Ashton
Chin, Jason W
Anderson, J Christopher
Schultz, Peter G

<120> UNNATURAL REACTIVE AMINO ACID GENETIC CODE ADDITIONS

<130> 54-000250US/PC

<160> 104

<170> PatentIn version 3.3

<210> 1
<211> 1275
<212> DNA
<213> Escherichia coli

<400> 1
atggcaagca gtaacttgat taaacaattg caagagcggg ggctggtagc ccaggtgacg 60

gacgaggaag cgttagcaga gcgactggcg caaggcccga tcgcgctcta ttgcggcttc 120

gatcctaccg ctgacagctt gcatttgggg catcttgttc cattgttatg cctgaaacgc 180

ttccagcagg cgggccacaa gccggttgcg ctggtaggcg gcgcgacggg tctgattggc 240

gacccgagct tcaaagctgc cgagcgtaag ctgaacaccg aagaaactgt tcaggagtgg 300

gtggacaaaa tccgtaagca ggttgccccg ttcctcgatt tcgactgtgg agaaaactct 360

gctatcgcgg cgaacaacta tgactggttc ggcaatatga atgtgctgac cttcctgcgc 420

gatattggca aacacttctc cgttaaccag atgatcaaca aagaagcggt taagcagcgt 480

ctcaaccgtg aagatcaggg gatttcgttc actgagtttt cctacaacct gttgcagggt 540

tatgacttcg cctgtctgaa caaacagtac ggtgtggtgc tgcaaattgg tggttctgac 600

cagtggggta acatcacttc tggtatcgac ctgacccgtc gtctgcatca gaatcaggtg 660

tttggcctga ccgttccgct gatcactaaa gcagatggca ccaaatttgg taaaactgaa 720

ggcggcgcag tctggttgga tccgaagaaa accagcccgt acaaattcta ccagttctgg 780

atcaacactg cggatgccga cgtttaccgc ttcctgaagt tcttcacctt tatgagcatt 840

gaagagatca acgccctgga agaagaagat aaaaacagcg gtaaagcacc gcgcgcccag 900

tatgtactgg cggagcaggt gactcgtctg gttcacggtg aagaaggttt acaggcggca 960

aaacgtatta ccgaatgcct gttcagcggt tctttgagtg cgctgagtga agcggacttc 1020

gaacagctgg cgcaggacgg cgtaccgatg gttgagatgg aaaagggcgc agacctgatg 1080

caggcactgg tcgattctga actgcaacct tcccgtggtc aggcacgtaa aactatcgcc 1140

tccaatgcca tcaccattaa cggtgaaaaa cagtccgatc ctgaatactt ctttaaagaa 1200

gaagatcgtc tgtttggtcg ttttaccttа ctgcgtcgcg gtaaaaagaa ttactgtctg 1260

atttgctgga aataa 1275

<210> 2
<211> 424
<212> PRT
<213> Escherichia coli

<400> 2

Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1 5 10 15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
20 25 30

Pro Ile Ala Leu Tyr Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
35 40 45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
85 90 95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
100 105 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
115 120 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
130 135 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145 150 155 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
165 170 175

Leu Leu Gln Gly Tyr Asp Phe Ala Cys Leu Asn Lys Gln Tyr Gly Val
180 185 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
195 200 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
210 215 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225 230 235 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
245 250 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
260 265 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
275 280 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
290 295 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305 310 315 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
325 330 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
340 345 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
355 360 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
370 375 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385 390 395 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
405 410 415

Asn Tyr Cys Leu Ile Cys Trp Lys
420

<210> 3
<211> 1275
<212> DNA
<213> Artificial

<220>
<223> artificial synthetase

<400> 3
atggcaagca gtaacttgat taaacaattg caagagcggg ggctggtagc ccaggtgacg 60

gacgaggaag cgttagcaga gcgactggcg caaggcccga tcgcactcgt gtgtggcttc 120

gatcctaccg ctgacagctt gcatttgggg catcttgttc cattgttatg cctgaaacgc 180

ttccagcagg cgggccacaa gccggttgcg ctggtaggcg gcgcgacggg tctgattggc 240

gacccgagct tcaaagctgc cgagcgtaag ctgaacaccg aagaaactgt tcaggagtgg 300

gtggacaaaa tccgtaagca ggttgccccg ttcctcgatt tcgactgtgg agaaaactct 360

gctatcgcgg ccaataatta tgactggttc ggcaatatga atgtgctgac cttcctgcgc 420

gatattggca aacacttctc cgttaaccag atgatcaaca aagaagcggt taagcagcgt 480

ctcaaccgtg aagatcaggg gatttcgttc actgagtttt cctacaacct gctgcagggt 540

tatagtatgg cctgtttgaa caaacagtac ggtgtggtgc tgcaaattgg tggttctgac 600

cagtggggta acatcacttc tggtatcgac ctgacccgtc gtctgcatca gaatcaggtg 660

tttggcctga ccgttccgct gatcactaaa gcagatggca ccaaatttgg taaaactgaa 720

ggcggcgcag tctggttgga tccgaagaaa accagcccgt acaaattcta ccagttctgg 780

atcaacactg cggatgccga cgtttaccgc ttcctgaagt tcttcacctt tatgagcatt 840

gaagagatca acgccctgga agaagaagat aaaaacagcg gtaaagcacc gcgcgcccag 900

tatgtactgg cggagcaggt gactcgtctg gttcacggtg aagaaggttt acaggcggca 960

aaacgtatta ccgaatgcct gttcagcggt tctttgagtg cgctgagtga agcggacttc 1020

gaacagctgg cgcaggacgg cgtaccgatg gttgagatgg aaaagggcgc agacctgatg 1080

caggcactgg tcgattctga actgcaacct tcccgtggtc aggcacgtaa aactatcgcc 1140

tccaatgcca tcaccattaa cggtgaaaaa cagtccgatc ctgaatactt ctttaaagaa 1200

gaagatcgtc tgtttggtcg ttttaccttа ctgcgtcgcg gtaaaaagaa ttactgtctg 1260

atttgctgga aataa 1275

<210> 4
<211> 1275
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 4
atggcaagca gtaacttgat taaacaattg caagagcggg ggctggtagc ccaggtgacg 60

gacgaggaag cgttagcaga gcgactggcg caaggcccga tcgcactcac ttgtggcttc 120

gatcctaccg ctgacagctt gcatttgggg catcttgttc cattgttatg cctgaaacgc 180

ttccagcagg cgggccacaa gccggttgcg ctggtaggcg gcgcgacggg tctgattggc 240

gacccgagct tcaaagctgc cgagcgtaag ctgaacaccg aagaaactgt tcaggagtgg 300

gtggacaaaa tccgtaagca ggttgccccg ttcctcgatt tcgactgtgg agaaaactct 360

gctatcgcgg ccaataatta tgactggttc agcaatatga atgtgctgac cttcctgcgc 420

gatattggca aacacttctc cgttaaccag atgatcaaca aagaagcggt taagcagcgt 480

ctcaaccgtg aagatcaggg gatttcgttc actgagtttt cctacaacct gctgcagggt 540

tatacgtatg cctgtctgaa caaacagtac ggtgtggtgc tgcaaattgg tggttctgac 600

cagtggggta acatcacttc tggtatcgac ctgacccgtc gtctgcatca gaatcaggtg 660

tttggcctga ccgttccgct gatcactaaa gcagatggca ccaaatttgg taaaactgaa 720

ggcggcgcag tctggttgga tccgaagaaa accagcccgt acaaattcta ccagttctgg 780

atcaacactg cggatgccga cgtttaccgc ttcctgaagt tcttcacctt tatgagcatt 840

gaagagatca acgccctgga agaagaagat aaaaacagcg gtaaagcacc gcgcgcccag 900

tatgtactgg cggagcaggt gactcgtctg gttcacggtg aagaaggttt acaggcggca 960

aaacgtatta ccgaatgcct gttcagcggt tctttgagtg cgctgagtga agcggacttc 1020

gaacagctgg cgcaggacgg cgtaccgatg gttgagatgg aaaagggcgc agacctgatg 1080

caggcactgg tcgattctga actgcaacct tcccgtggtc aggcacgtaa aactatcgcc 1140

tccaatgcca tcaccattaa cggtgaaaaa cagtccgatc ctgaatactt ctttaaagaa 1200

gaagatcgtc tgtttggtcg ttttaccttа ctgcgtcgcg gtaaaaagaa ttactgtctg 1260

atttgctgga aataa 1275

<210> 5
<211> 1275
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 5
atggcaagca gtaacttgat taaacaattg caagagcggg ggctggtagc ccaggtgacg 60

gacgaggaag cgttagcaga gcgactggcg caaggcccga tcgcactcgt gtgtggcttc 120

gatcctaccg ctgacagctt gcatttgggg catcttgttc cattgttatg cctgaaacgc 180

ttccagcagg cgggccacaa gccggttgcg ctggtaggcg gcgcgacggg tctgattggc 240

gacccgagct tcaaagctgc cgagcgtaag ctgaacaccg aagaaactgt tcaggagtgg 300

gtggacaaaa tccgtaagca ggttgccccg ttcctcgatt tcgactgtgg agaaaactct 360

gctatcgcgg ccaataatta tgactggttc ggcaatatga atgtgctgac cttcctgcgc 420

gatattggca aacacttctc cgttaaccag atgatcaaca aagaagcggt taagcagcgt 480

ctcaaccgtg aagatcaggg gatttcgttc actgagtttt cctacaacct gctgcagggt 540

tatagtatgg cctgtttgaa caaacagtac ggtgtggtgc tgcaaattgg tggttctgac 600

cagtggggta acatcacttc tggtatcgac ctgacccgtc gtctgcatca gaatcaggtg 660

tttggcctga ccgttccgct gatcactaaa gcagatggca ccaaatttgg taaaactgaa 720

ggcggcgcag tctggttgga tccgaagaaa accagcccgt acaaattcta ccagttctgg 780

atcaacactg cggatgccga cgtttaccgc ttcctgaagt tcttcacctt tatgagcatt 840

gaagagatca acgccctgga agaagaagat aaaaacagcg gtaaagcacc gcgcgcccag 900

tatgtactgg cggagcaggt gactcgtctg gttcacggtg aagaaggttt acaggcggca 960

aaacgtatta ccgaatgcct gttcagcggt tctttgagtg cgctgagtga agcggacttc 1020

gaacagctgg cgcaggacgg cgtaccgatg gttgagatgg aaaagggcgc agacctgatg 1080

caggcactgg tcgattctga actgcaacct tcccgtggtc aggcacgtaa aactatcgcc 1140

tccaatgcca tcaccattaa cggtgaaaaa cagtccgatc ctgaatactt ctttaaagaa 1200

gaagatcgtc tgtttggtcg ttttaccttа ctgcgtcgcg gtaaaaagaa ttactgtctg 1260

atttgctgga aataa 1275

<210> 6
<211> 1275
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 6
atggcaagca gtaacttgat taaacaattg caagagcggg ggctggtagc ccaggtgacg 60

gacgaggaag cgttagcaga gcgactggcg caaggcccga tcgcactcgt gtgtggcttc 120

gatcctaccg ctgacagctt gcatttgggg catcttgttc cattgttatg cctgaaacgc 180

ttccagcagg cgggccacaa gccggttgcg ctggtaggcg gcgcgacggg tctgattggc 240

gacccgagct tcaaagctgc cgagcgtaag ctgaacaccg aagaaactgt tcaggagtgg 300

gtggacaaaa tccgtaagca ggttgccccg ttcctcgatt tcgactgtgg agaaaactct 360

gctatcgcgg ccaataatta tgactggttc ggcaatatga atgtgctgac cttcctgcgc 420

gatattggca aacacttctc cgttaaccag atgatcaaca aagaagcggt taagcagcgt 480

ctcaaccgtg aagatcaggg gatttcgttc actgagtttt cctacaacct gctgcagggt 540

tatagtatgg cctgtttgaa caaacagtac ggtgtggtgc tgcaaattgg tggttctgac 600

cagtggggta acatcacttc tggtatcgac ctgacccgtc gtctgcatca gaatcaggtg 660

tttggcctga ccgttccgct gatcactaaa gcagatggca ccaaatttgg taaaactgaa 720

ggcggcgcag tctggttgga tccgaagaaa accagcccgt acaaattcta ccagttctgg 780

atcaacactg cggatgccga cgtttaccgc ttcctgaagt tcttcacctt tatgagcatt 840

gaagagatca acgccctgga agaagaagat aaaaacagcg gtaaagcacc gcgcgcccag 900

tatgtactgg cggagcaggt gactcgtctg gttcacggtg aagaaggttt acaggcggca 960

aaacgtatta ccgaatgcct gttcagcggt tctttgagtg cgctgagtga agcggacttc 1020

gaacagctgg cgcaggacgg cgtaccgatg gttgagatgg aaaagggcgc agacctgatg 1080

caggcactgg tcgattctga actgcaacct tcccgtggtc aggcacgtaa aactatcgcc 1140

tccaatgcca tcaccattaa cggtgaaaaa cagtccgatc ctgaatactt ctttaaagaa 1200

gaagatcgtc tgtttggtcg ttttaccttа ctgcgtcgcg gtaaaaagaa ttactgtctg 1260

atttgctgga aataa 1275

<210> 7
<211> 1275
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 7
atggcaagca gtaacttgat taaacaattg caagagcggg ggctggtagc ccaggtgacg 60

gacgaggaag cgttagcaga gcgactggcg caaggcccga tcgcactcac gtgtggcttc 120

gatcctaccg ctgacagctt gcatttgggg catcttgttc cattgttatg cctgaaacgc 180

ttccagcagg cgggccacaa gccggttgcg ctggtaggcg gcgcgacggg tctgattggc 240

gacccgagct tcaaagctgc cgagcgtaag ctgaacaccg aagaaactgt tcaggagtgg 300

gtggacaaaa tccgtaagca ggttgccccg ttcctcgatt tcgactgtgg agaaaactct 360

gctatcgcgg ccaataatta tgactggttc ggcaatatga atgtgctgac cttcctgcgc 420

gatattggca aacacttctc cgttaaccag atgatcaaca aagaagcggt taagcagcgt 480

ctcaaccgtg aagatcaggg gatttcgttc actgagtttt cctacagcct gctgcagggt 540

tatacgatgg cctgtctgaa caaacagtac ggtgtggtgc tgcaaattgg tggttctgac 600

cagtggggta acatcacttc tggtatcgac ctgacccgtc gtctgcatca gaatcaggtg 660

tttggcctga ccgttccgct gatcactaaa gcagatggca ccaaatttgg taaaactgaa 720

ggcggcgcag tctggttgga tccgaagaaa accagcccgt acaaattcta ccagttctgg 780

atcaacactg cggatgccga cgtttaccgc ttcctgaagt tcttcacctt tatgagcatt 840

gaagagatca acgccctgga agaagaagat aaaaacagcg gtaaagcacc gcgcgcccag 900

tatgtactgg cggagcaggt gactcgtctg gttcacggtg aagaaggttt acaggcggca 960

aaacgtatta ccgaatgcct gttcagcggt tctttgagtg cgctgagtga agcggacttc 1020

gaacagctgg cgcaggacgg cgtaccgatg gttgagatgg aaaagggcgc agacctgatg 1080

caggcactgg tcgattctga actgcaacct tcccgtggtc aggcacgtaa aactatcgcc 1140

tccaatgcca tcaccattaa cggtgaaaaa cagtccgatc ctgaatactt ctttaaagaa 1200

gaagatcgtc tgtttggtcg ttttaccttа ctgcgtcgcg gtaaaaagaa ttactgtctg 1260

atttgctgga aataa 1275

<210> 8
<211> 540

<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 8
cgggggctgg tagcccaggt gacggacgag gaagcgttag cagagcgact ggcgcaaggc 60

ccgatcgcac tcacttgtgg cttcgatcct accgctgaca gcttgcattt ggggcatctt 120

gttccattgt tatgcctgaa acgcttccag caggcgggcc acaagccggt tgcgctggta 180

ggcggcgcga cgggtctgat tggcgacccg agcttcaaag ctgccgagcg taagctgaac 240

accgaagaaa ctgttcagga gtgggtggac aaaatccgta agcaggttgc cccgttcctc 300

gatttcgact gtggagaaaa ctctgctatc gcggccaata ttatgactg gttcagcaat 360

atgaatgtgc tgaccttcct gcgcgatatt ggcaaacact tctccgttaa ccagatgatc 420

aacaaagaag cggttaagca gcgtctcaac cgtgaagatc aggggatttc gttcactgag 480

ttttcctaca acctgctgca gggttatacg tatgcctgtc tgaacaaaca gtacggtgtg 540

<210> 9
<211> 540
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 9
cgggggctgg taccccaggt gacggacgag gaagcgttag cagagcgact ggcgcaaggc 60

ccgatcgcac tcacttgtgg cttcgatcct accgctgaca gcttgcattt ggggcatctt 120

gttccattgt tatgcctgaa acgcttccag caggcgggcc acaagccggt tgcgctggta 180

ggcggcgcga cgggtctgat tggcgacccg agcttcaaag ctgccgagcg taagctgaac 240

accgaagaaa ctgttcagga gtgggtggac aaaatccgta agcaggttgc cccgttcctc 300

gatttcgact gtggagaaaa ctctgctatc gcggccaata ttatgactg gttcagcaat 360

atgaatgtgc tgaccttcct gcgcgatatt ggcaaacact tctccgttaa ccagatgatc 420

aacaaagaag cggttaagca gcgtctcaac cgtgaagatc aggggatttc gttcactgag 480

ttttcctaca acctgctgca gggttatacg tatgcctgtc tgaacaaaca gtacggtgtg 540

<210> 10
<211> 540
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 10
cgggggctgg tagcccaggt gacggacgag gaagcgttag cagagcgact ggcgcaaggc 60

ccgatcgcac tcacttgtgg cttcgatcct accgctgaca gcttgcattt ggggcatctt 120

gttccattgt tatgcctgaa acgcttccag caggcgggcc acaagccggt tgcgctggta 180

ggcggcgcga cgggtctgat tggcgacccg agcttcaaag ctgccgagcg taagctgaac 240

accgaagaaa ctgttcagga gtgggtggac aaaatccgta agcaggttgc cccgttcctc 300

gatttcgact gtggagaaaa ctctgctatc gcggccaata attatgactg gttcagcaat 360

atgaatgtgc tgaccttcct gcgcgatatt ggcaaacact tctccgttaa ccagatgatc 420

aacaaagaag cggttaagca gcgtctcaac cgtgaagatc aggggatttc gttcactgag 480

ttttcctaca acctgctgca gggttatacg tatgcctgtc tgaacaaaca gtacggtgtg 540

<210> 11
<211> 540
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 11
cgggggctgg tagcccaggt gacggacgag gaagcgttag cagagcgact ggcgcaaggc 60

ccgatcgcac tcacttgtgg cttcgatcct accgctgaca gcttgcattt ggggcatctt 120

gttccattgt tatgcctgaa acgcttccag caggcgggcc acaagccggt tgcgctggta 180

ggcggcgcga cgggtctgat tggcgacccg agcttcaaag ctgccgagcg taagctgaac 240

accgaagaaa ctgttcagga gtgggtggac aaaatccgta agcaggttgc cccgttcctc 300

gatttcgact gtggagaaaa ctctgctatc gcggccaata attatgactg gttcggcaat 360

atgaatgtgc tgaccttcct gcgcgatatt ggcaaacact tctccgttaa ccagatgatc 420

aacaaagaag cggttaagca gcgtctcaac cgtgaagatc aggggatttc gttcactgag 480

ttttcctaca acctgctgca gggttattcg tatgcctgtg cgaacaaaca gtacggtgtg 540

<210> 12
<211> 540
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 12
cgggggctgg tagcccaggt gacggacgag gaagcgttag cagagcgact ggcgcaaggc 60

ccgatcgcac tcacttgtgg cttcgatcct accgctgaca gcttgcattt ggggcatctt 120

gttccattgt tatgcctgaa acgcttccag caggcgggcc acaagccggt tgcgctggta 180

ggcggcgcga cgggtctgat tggcgacccg agcttcaaag ctgccgagcg taagctgaac 240

accgaagaaa ctgttcagga gtgggtggac aaaatccgta agcaggttgc cccgttcctc 300

gatttcgact gtggagaaaa ctctgctatc gcggccaata attatgactg gttcagcaat 360

atgaatgtgc tgaccttcct gcgcgatatt ggcaaacact tctccgttaa ccagatgatc 420

aacaaagaag cggttaagca gcgtctcaac cgtgaagatc aggggatttc gttcactgag 480

ttttcctaca acctgctgca gggttatacg tatgcctgtc tgaacaaaca gtacggtgtg 540

<210> 13
<211> 540
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 13
cgggggctgg taccccaggt gacggacgag gaagcgttag cagagcgact ggcgcaaggc 60

ccgatcgcac tcctttgtgg cttcgatcct accgctgaca gcttgcattt ggggcatctt 120

gttccattgt tatgcctgaa acgcttccag caggcgggcc acaagccggt tgcgctggta 180

ggcggcgcga cgggtctgat tggcgacccg agcttcaaag ctgccgagcg taagctgaac 240

accgaagaaa ctgttcagga gtgggtggac aaaatccgta agcaggttgc cccgttcctc 300

gatttcgact gtggagaaaa ctctgctatc gcggccaata ttatgactg gttcggcaat 360

atgaatgtgc tgaccttcct gcgcgatatt ggcaaacact tctccgttaa ccagatgatc 420

aacaaagaag cggttaagca gcgtctcaac cgtgaagatc aggggatttc gttcactgag 480

ttttcctaca acctgctgca gggttattct attgcctgtt cgaacaaaca gtacggtgtg 540

<210> 14
<211> 540
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 14
cgggggctgg tagcccaggt gacggacgag gaagcgttag cagagcgact ggcgcaaggc 60

ccgatcgcac tcgtgtgtgg cttcgatcct accgctgaca gcttgcattt ggggcatctt 120

gttccattgt tatgcctgaa acgcttccag caggcgggcc acaagccggt tgcgctggta 180

ggcggcgcga cgggtctgat tggcgacccg agcttcaaag ctgccgagcg taagctgaac 240

accgaagaaa ctgttcagga gtgggtggac aaaatccgta agcaggttgc cccgttcctc 300

gatttcgact gtggagaaaa ctctgctatc gcggccaata ttatgactg gttcggcaat 360

atgaatgtgc tgaccttcct gcgcgatatt ggcaaacact tctccgttaa ccagatgatc 420

aacaaagaag cggttaagca gcgtctcaac cgtgaagatc aggggatttc gttcactgag 480

ttttcctaca acctgctgca gggttatagt attgcctgtt tgaacaaaca gtacggtgtg 540

<210> 15

<211> 540
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 15
cgggggctgg taccccaggt gacggacgag gaagcgttag cagagcgact ggcgcaaggc 60

ccgatcgcac tcgtgtgtgg cttcgatcct accgctgaca gcttgcattt ggggcatctt 120

gttccattgt tatgcctgaa acgcttccag caggcgggcc acaagccggt tgcgctggta 180

ggcggcgcga cgggtctgat tggcgacccg agcttcaaag ctgccgagcg taagctgaac 240

accgaagaaa ctgttcagga gtgggtggac aaaatccgta agcaggttgc cccgttcctc 300

gatttcgact gtggagaaaa ctctgctatc gcggccaata attatgactg gttcggcaat 360

atgaatgtgc tgaccttcct gcgcgatatt ggcaaacact tctccgttaa ccagatgatc 420

aacaaagaag cggttaagca gcgtctcaac cgtgaagatc aggggatttc gttcactgag 480

ttttcctaca acctgctgca gggttatagt attgcctgtt tgaacaaaca gtacggtgtg 540

<210> 16
<211> 540
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 16
cgggggctgg tagcccaggt gacggacgag gaagcgttag cagagcgact ggcgcaaggc 60

ccgatcgcac tctggtgtgg cttcgatcct accgctgaca gcttgcattt ggggcatctt 120

gttccattgt tatgcctgaa acgcttccag caggcgggcc acaagccggt tgcgctggta 180

ggcggcgcga cgggtctgat tggcgacccg agcttcaagg ctgccgagcg taagctgaac 240

accgaagaaa ctgttcagga gtgggtggac aaaatccgta agcaggttgc cccgttcctc 300

gatttcgact gtggagaaaa ctctgctatc gcggccaatt gttatgactg gttcggcaat 360

atgaatgtgc tgaccttcct gcgcgatatt ggcaaacact tctccgttaa ccagatgatc 420

aacaaagaag cggttaagca gcgtctcaac cgtgaagatc aggggatttc gttcactgag 480

ttttcctaca acctgctgca gggttatatg cgtgcctgtg agaacaaaca gtacggtgtg 540

<210> 17
<211> 624
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 17
cgggggctgg tagcccaggt gacggacgag gaagcgttag cagagcgact ggcgcaaggc 60

ccgatcgcac tcatttgtgg cttcgatcct accgctgaca gcttgcattt ggggcatctt 120

gttccattgt tatgcctgaa acgcttccag caggcgggcc acaagccggt tgcgctggta 180

ggcggcgcga cgggtctgat tggcgacccg agcttcaaag ctgccgagcg taagctgaac 240

accgaagaaa ctgttcagga gtgggtggac aaaatccgta agcaggttgc cccgttcctc 300

gatttcgact gtggagaaaa ctctgctatc gcggccaata attatgactg gttcggcaat 360

atgaatgtgc tgaccttcct gcgcgatatt ggcaaacact tctccgttaa ccagatgatc 420

aacaaagaag cggttaagca gcgtctcaac cgtgaaggtc aggggatttc gttcactgag 480

ttttcctaca acctgctgca gggttatggt atggcctgtg ctaacaaaca gtacggtgtg 540

gtgctgcaaa ttggtggttc tgaccaatgg ggtaacatca cttctggtat cgacctgacc 600

cgtcgtctgc atcagaatca ggtg 624

<210> 18
<211> 609
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 18
caggtgacgg acgaggaagc gttagcagag cgactggcgc aaggcccgat cgcactcggt 60

tgtggcttcg atcctaccgc tgacagcttg catttggggc atcttgttcc attgttatgc 120

ctgaaacgct tccagcaggc gggccacaag ccggttgcgc tggtaggcgg cgcgacgggt 180

ctgattggcg acccgagctt caaagctgcc gagcgtaagc tgaacaccga agaaactgtt 240

caggagtggg tggacaaaat ccgtaagcag gttgccccgt tcctcgattt cgactgtgga 300

gaaaactctg ctatcgcggc caataattat gactggttcg gcaatatgaa tgtgctgacc 360

ttcctgcgcg atattggcaa acacttctcc gttaaccaga tgatcaacaa agaagcggtt 420

aagcagcgtc tcaaccgtga agatcagggg atttcgttca ctgagttttc ctacaacctg 480

ctgcagggtt atggttttgc ctgtttgaac aaacagtacg gtgtggtgct gcaaattggt 540

ggttctgacc agtggggtaa catcacttct ggtatcgacc tgacccgtcg tctgcatcag 600

aatcaggtg 609

<210> 19
<211> 591
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 19
gcgttagcag agcgactggc gcaaggcccg atcgcactcg ggtgtggctt cgatcctacc 60

gctgacagct tgcatttggg gcatcttgtt ccattgttat gcctgaaacg cttccagcag 120

gcgggccaca agccggttgc gctggtaggc ggcgcgacgg gtctgattgg cgacccgagc 180

ttcaaagctg ccgagcgtaa gctgaacacc gaagaaactg ttcaggagtg ggtggacaaa 240

atccgtaagc aggttgcccc gttcctcgat ttcgactgtg gagaaaactc tgctatcgcg 300

gccaataatt atgactggtt cggcaatatg aatgtgctga ccttcctgcg cgatattggc 360

aaacacttct ccgttaacca gatgatcaac aaagaagcgg ttaagcagcg tctcaaccgt 420

gaagatcagg ggatttcgtt cactgagttt tcctacaacc tgctgcaggg ttatggttat 480

gcctgtatga acaaacagta cggtgtggtg ctgcaaattg gtggttctga ccagtggggt 540

aacatcactt ctggtatcga cctgacccgt cgtctgcatc agaatcaggt g 591

<210> 20
<211> 621
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<220>
<221> misc_feature
<222> (26)..(26)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (612)..(612)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (618)..(618)
<223> n is a, c, g, or t

<400> 20
gggctggtag cccaggtgac ggacgnagaa gcgttagcag agcgactggc gcaaggcccg 60

atcgcactcc tttgtggctt cgatcctacc gctgacagct tgcatttggg gcatcttgtt 120

ccattgttat gcctgaaacg cttccagcag gcgggccaca agccggttgc gctggtaggc 180

ggcgcgacgg gtctgattgg cgacccgagc ttcaaagctg ccgagcgtaa gctgaacacc 240

gaagaaactg ttcaggagtg ggtggacaaa atccgtaagc aggttgcccc gttcctcgat 300

ttcgactgtg gagaaaactc tgctatcgcg gccaataatt atgactggtt cggcaatatg 360

aatgtgctga ccttcctgcg cgatattggc aaacacttct ccgttaacca gatgatcaac 420

aaagaagcgg ttaagcagcg tctcaaccgt gaagatcagg ggatttcgtt cactgagttt 480

tcctacaacc tgctgcaggg ttattctatg gcctgtgcga acaaacagta cggtgtggtg 540

ctgcaaattg gtggttctga ccagtggggt aacatcactt ctggtatcga cctgacccgt 600

cgtctgcatc anaatcangt g 621

<210> 21
<211> 588
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 21
ttagcagagc gactggcgca aggcccgatc gcactcgttt gtggcttcga tcctaccgct 60

gacagcttgc atttggggca tcttgttcca ttgttatgcc tgaaacgctt ccagcaggcg 120

ggccacaagc cggttgcgct ggtaggcggc gcgacgggtc tgattggcga cccgagcttc 180

aaagctgccg agcgtaagct gaacaccgaa gaaactgttc aggagtgggt ggacaaaatc 240

cgtaagcagg ttgccccgtt cctcgatttc gactgtggag aaaactctgc tatcgcggcc 300

aataattatg actggttcgg caatatgaat gtgctgacct tcctgcgcga tattggcaaa 360

cacttctccg ttaaccagat gatcaacaaa gaagcggtta agcagcgtct caaccgtgaa 420

gatcagggga tttcgttcac tgagttttcc tacaacctgc tgcagggtta ttctgcggcc 480

tgtgcgaaca aacagtacgg tgtggtgctg caaattggtg gttctgacca gtggggtaac 540

atcacttctg gtatcgacct gacccgtcgt ctgcatcaga atcaggtg 588

<210> 22
<211> 600
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<220>
<221> misc_feature
<222> (403)..(403)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (513)..(513)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (515)..(515)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (518)..(518)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (531)..(531)
<223> n is a, c, g, or t

<400> 22
gacgaggaag cgttagcaga gcgactggcg caaggcccga tcgcactcct gtgtggcttc 60

gatcctaccg ctgacagctt gcatttgggg catcttgttc cattgttatg cctgaaacgc 120

ttccagcagg cgggccacaa gccggttgcg ctggtaggcg gcgcgacggg tctgattggc 180

gacccgagct tcaaagctgc cgagcgtaag ctgaacaccg aagaaactgt tcaggagtgg 240

gtggacaaaa tccgtaagca ggttgccccg ttcctcgatt tcgactgtgg agaaaactct 300

gctatcgcgg ccaataatta tgactggttc ggcaatatga atgtgctgac cttcctgcgc 360

gatattggca aacacttctc cgttaaccag atgatcaaca aanaagcggt taagcagcgt 420

ctcaaccgtg aagatcaggg gatttcgttc actgagtttt cctacaacct gctgcagggt 480

tattcggctg cctgtgcgaa caaacagtac ggngnggngc tgcaaattgg nggttctgac 540

caggggggta acatcacttc tggtatcgac ctgacccgtc gtctgcatca aaatcaggtg 600

<210> 23
<211> 591
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<220>
<221> misc_feature
<222> (588)..(588)
<223> n is a, c, g, or t

<400> 23
gcgttagcag agcgactggc gcaaggcccg atcgcactcg tttgtggctt cgatcctacc 60

gctgacagct tgcatttggg gcatcttgtt ccattgttgt gcctgaaacg cttccagcag 120

gcgggccaca agccggttgc gctggtaggc ggcgcgacgg gtctgattgg cgacccgagc 180

ttcaaagctg ccgagcgtaa gctgaacacc gaagaaactg ttcaggagtg ggtggacaaa 240

atccgtaagc aggttgcccc gttcctcgat ttcgactgtg gagaaaactc tgctatcgcg 300

gccaataatt atgactggtt cggcaatatg aatgtgctga ccttcctgcg cgatattggc 360

aaacacttct ccgttaacca gatgatcaac aaagaagcgg ttaagcagcg tctcaaccgt 420

gaagatcagg ggatttcgtt cactgagttt tcctacaacc tgctgcaggg ttatagtgcg 480

gcctgtgtta acaaacagta cggtgtggtg ctgcaaattg gtggttctga ccagtggggt 540

aacatcactt ctggtatcga cctgacccgt cgtctgcatc agaatcangt g 591

<210> 24
<211> 600
<212> DNA
<213> artificial

<220>

<223> artificial synthetase

<400> 24
gacgaggaag cgttagcaga gcgactggcg caaggcccga tcgcactcat ttgtggcttc 60

gatcctaccg ctgacagctt gcatttgggg catcttgttc cattgttatg cctgaaacgc 120

ttccagcagg cgggccacaa gccggttgcg ctggtaggcg gcgcgacggg tctgattggc 180

gacccgagct tcaaagctgc cgagcgtaag ctgaacaccg aagaaactgt tcaggagtgg 240

gtggacaaaa tccgtaagca ggttgccccg ttcctcgatt tcgactgtgg agaaaactct 300

gctatcgcgg ccaatgatta tgactggttc ggcaatatga atgtgctgac cttcctgcgc 360

gatattggca aacacttctc cgttaaccag atgatcaaca aagaagcggt taagcagcgt 420

ctcaaccgtg aagatcaggg gatttcgttc actgagtttt cctacaacct gctgcagggt 480

tataattttg cctgtgtgaa caaacagtac ggtgtggtgc tgcaaattgg tggttctgac 540

cagtggggta acatcacttc tggtatcgac ctgacccgtc gtctgcatca gaatcaggtg 600

<210> 25
<211> 579
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 25
cgactggcgc aaggcccgat cgcactcacg tgtggcttcg atcctaccgc tgacagcttg 60

catttggggc atcttgttcc attgttatgc ctgaaacgct tccagcaggc gggccacaag 120

ccggttgcgc tggtaggcgg cgcgacgggt ctgattggcg acccgagctt caaagctgcc 180

gagcgtaagc tgaacaccga agaaactgtt caggagtggg tggacaaaat ccgtaagcag 240

gttgccccgt tcctcgattt cgactgtgga gaaaactctg ctatcgcggc caataattat 300

gactggttcg gcaatatgaa tgtgctgacc ttcctgcgcg atattggcaa acacttctcc 360

gttaaccaga tgatcaacaa agaagcggtt aagcagcgtc tcaaccgtga agatcagggg 420

atttcgttca ctgagttttc ctacaatctg ctgcagggtt attcggctgc ctgtcttaac 480

aaacagtacg gtgtggtgct gcaaattggt ggttctgacc agtggggtaa catcacttct 540

ggtatcgacc tgacccgtcg tctgcatcag aatcaggtg 579

<210> 26
<211> 624
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<220>
<221> misc_feature

<222> (13)..(13)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (599)..(599)
<223> n is a, c, g, or t

<400> 26
cgggggctgg tancccaggt gacggacgag gaagcgttag cagagcgact ggcgcaaggc 60

ccgatcgcac tcgggtgtgg cttcgatcct accgctgaca gcttgcattt ggggcatctt 120

gttccattgt tatgcctgaa acgcttccag caggcgggcc acaagccggt tgcgctggta 180

ggcggcgcga cgggtctgat tggcgacccg agcttcaaag ctgccgagcg taagctgaac 240

accgaagaaa ctgttcagga gtgggtggac aaaatccgta agcaggttgc cccgttcctc 300

gatttcgact gtggagaaaa ctctgctatc gcggccaata attatgactg gttcggcaat 360

atgaatgtgc tgaccttcct gcgcgatatt ggcaaacact tctccgttaa ccagatgatc 420

aacaaagaag cggttaagca gcgtctcaac cgtgaagatc aggggatttc gttcactgag 480

ttttcctaca acctgctgca gggttattct atggcctgtt tgaacaaaca gtacggtgtg 540

gtgctgcaaa ttggtggttc tgaccagtgg ggtaacatca cttctggtat cgacctganc 600

cgtcgtctgc atcagaatca ggtg 624

<210> 27
<211> 625
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<220>
<221> misc_feature
<222> (600)..(600)
<223> n is a, c, g, or t

<400> 27
cgggggctgg tagcccaggt gacggacgag gaagcgttag cagagcgact ggcgcaaggc 60

ccgatcgcac tcacgtgtgg cttcgatcct accgctgaca gcttgcattt ggggcatctt 120

gttccattgt tatgcctgaa acgcttccag caggcgggcc acaagccggt tgcgctggta 180

ggcggcgcga cgggtctgat tggcgacccg agcttcaaag ctgccgagcg taagctgaac 240

accgaagaaa ctgttcagga gtgggtggac aaaatccgta agcaggttgc cccgttcctc 300

gatttcgact gtggagaaaa ctctgctatc gcggccaata attatgactg gttcggcaat 360

atgaatgtgc tgaccttcct gcgcgatatt ggcaaacact tctccgttaa ccagatgatc 420

aacaaagaag cggttaagca gcgtctcaac cgtgaagatc aggggatttc gttcactgag 480

ttttcctaca atctgctgca gggttattcg gctgcctgtc ttaacaaaca gtacggtgtg 540

gtgctgcaaa ttggtggttc tgaccagtgg ggtaacatca cttctggtat cgaacctgan 600

ccgtcgtctg catcaaaatc aagtg 625

<210> 28
<211> 624
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 28
cgggggctgg taccccaagt gacggacgag gaaacgttag cagagcgact ggcgcaaggc 60

ccgatcgcac tctcttgtgg cttcgatcct accgctgaca gcttgcattt ggggcatctt 120

gttccattgt tatgcctgaa acgcttccag caggcaggcc acaagccggt tgcgctggta 180

ggcggcgcga cgggtctgat tggcgacccg agcttcaaag ctgccgagcg taagctgaac 240

accgaagaaa ctgttcagga gtgggtggac aaaatccgta agcaggttgc cccgttcctc 300

gatttcgact gtggagaaaa ctctgctatc gcggccaata attatgactg gttcggcaat 360

atgaatgtgc tgaccttcct gcgcgatatt ggcaaacact tctccgttaa ccagatgatc 420

aacaaagaag cggttaagca gcgtctcaac cgtgaagatc aggggatttc gttcactgag 480

ttttcctaca acctgctgca gggttatacg atggcctgtg tgaacaaaca gtacggtgtg 540

gtgctgcaaa ttggtggttc tgaccagtgg ggtaacatca cttctggtat cgacctgacc 600

cgtcgtctgc atcagaatca ggtg 624

<210> 29
<211> 624
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 29
cgggggctgg tagcccaggt gacggacgag gaagcgttag cagagcgact ggcgcaaggc 60

ccgatcgcac tcgcgtgcgg cttcgatcct accgctgaca gcttgcattt ggggcatctt 120

gttccattgt tatgcctgaa acgcttccag caggcgggcc acaagccggt tgcgctggta 180

ggcggcgcga cgggtctgat tggcgacccg agcttcaagg ctgccgagcg taagctgaac 240

accgaagaaa ctgttcagga gtgggtggac aaaatccgta agcaggttgc cccgttcctc 300

gatttcgact gtggagaaaa ctctgctatc gcggccaata attatgactg gttcggcaat 360

atgaatgtgc tgaccttcct gcgcgatatt ggcaaacact tctccgttaa ccagatgatc 420

aacaaagaag cggttaagca gcgtctcaac cgtgaagatc aggggatttc gttcactgag 480

ttttcctaca acctgctgca gggttattct tatgcctgtc ttaacaaaca gtacggtgtg 540

gtgctgcaaa ttggtggttc tgaccagtgg ggtaacatca cttctggtat cgacctgacc 600

cgtcgtctgc atcagaatca ggtg 624

<210> 30
<211> 624
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 30
cgggggctgg tagcccaggt gacggacgag gaagcgttag cagagcgact ggcgcaaggc 60

ccgatcgcac tcgcgtgtgg cttcgatcct accgctgaca gcttgcattt ggggcatctt 120

gttccattgt tatgcctgaa acgcttccag caggcgggcc acaagccggt tgcgctggta 180

ggcggcgcga cgggtctgat tggcgacccg agcttcaaag ctgccgagcg taagctgaac 240

accgaagaaa ctgttcagga gtgggtggac aaaatccgta agcaggttgc cccgttcctc 300

gatttcgact gtggagaaaa ctctgctatc gcggccaata attatgactg gttcggcaat 360

atgaatgtgc tgaccttcct gcgcgatatt ggcaaacact tctccgttaa ccagatgatc 420

aacaaagaag cggttaagca gcgtctcaac cgtgaagatc aggggatttc gttcactgag 480

ttttcctaca acctgctgca gggttatacg atggcctgtt gtaacaaaca gtacggtgtg 540

gtgctgcaaa ttggtggttc tgaccagtgg ggtaacatca cttctggtat cgacctgacc 600

cgtcgtctgc atcagaatca ggtg 624

<210> 31
<211> 624
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 31
cgggggctgg taccccaagt gacggacgag gaagcgttag cagagcgact ggcgcaaggc 60

ccgatcgcac tcacgtgtgg cttcgatcct accgctgaca gcttgcattt ggggcatctt 120

gttccattgt tatgcctgaa acgcttccag caggcgggcc acaagccggt tgcgctggta 180

ggcggcgcga cgggtctgat tggcgacccg agcttcaaag ctgccgagcg taagctgaac 240

accgaagaaa ctgttcagga gtgggtggac aaaatccgta agcaggttgc cccgttcctc 300

gatttcgact gtggagaaaa ctctgctatc gcggccaata attatgactg gttcggcaat 360

atgaatgtgc tgaccttcct gcgcgatatt ggcaaacact tctccgttaa ccagatgatc 420

aacaaagaag cggttaagca gcgtctcaac cgtgaagatc aggggatttc gttcgctgag 480

ttttcctaca acctgctgca gggttatacg tttgcctgta tgaacaaaca gtacggtgtg 540

gtgctgcaaa ttggtggttc tgaccagtgg ggtaacatca cttctggtat cgacctgacc 600

cgtcgtctgc atcagaatca ggtg 624

<210> 32
<211> 606
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 32
gtgacggacg aggaagcgtt agcagagcga ctggcgcaag gcccgatcgc actcacgtgt 60

ggcttcgatc ctaccgctga cagcttgcat ttggggcatc ttgttccatt gttatgcctg 120

aaacgcttcc agcaggcggg ccacaagccg gttgcgctgg taggcggcgc gacgggtctg 180

attggcgacc cgagcttcaa agctgccgag cgtaagctga acaccgaaga aactgttcag 240

gagtgggtgg acaaaatccg taagcaggtt gccccgttcc tcgatttcga ctgtggagaa 300

aactctgcta tcgcggccaa taattatgac tggttcggca atatgaatgt gctgaccttc 360

ctgcgcgata ttggcaaaca cttctccgtt aaccagatga tcaacaaaga agcggttaag 420

cagcgtctca accgtgaaga tcaggggatt tcgttcactg agttttccta caatctgctg 480

cagggttatt cggctgcctg tcttaacaaa cagtacggtg tggtgctgca aattggtggt 540

tctgaccagt ggggtaacat cacttctggt atcgacctga cccgtcgtct gcatcagaat 600

caggtg 606

<210> 33
<211> 624
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 33
cgggggctgg tagcccaggt gacggacgag gaagcgttag cagagcgact ggcgcaaggc 60

ccgatcgcac tcgtttgtgg cttcgatcct accgctgaca gcttgcattt ggggcatctt 120

gttccattgt tatgcctgaa acgcttccag caggcgggcc acaagccggt tgcgctggta 180

ggcggcgcga cgggtctgat tggcgacccg agcttcaaag ctgccgagcg taagctgaac 240

accgaagaaa ctgttcagga gtgggtggac aaaatccgta agcaggttgc cccgttcctc 300

gatttcgact gtggagaaaa ctctgctatc gcggccaata attatgactg gttcggcaat 360

atgaatgtgc tgaccttcct gcgcgatatt ggcaaacact tctccgttaa ccagatgatc 420

aacaaagaag cggttaagca gcgtctcaac cgtgaagatc aggggatttc gttcactgag 480

ttttcctaca acctgctgca gggttattcg atggcctgta cgaacaaaca gtacggtgtg 540

gtgctgcaaa ttggtggttc tgaccagtgg ggtaacatca cttctggtat cgacctgacc 600

cgtcgtctgc atcagaatca ggtg 624

<210> 34
<211> 624
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<220>
<221> misc_feature
<222> (13)..(13)
<223> n is a, c, g, or t

<400> 34
cgggggctgg tancccaagt gacggacggg gaagcgttag cagagcgact ggcgcaaggc 60

ccgatcgcac tcagttgtgg cttcgatcct accgctgaca gcttgcattt ggggcatctt 120

gttccattgt tatgcctgaa acgcttccag caggcgggcc acaagccggt tgcgctggta 180

ggcggcgcga cgggtctgat tggcgacccg agcttcaaag ctgccgagcg taagctgaac 240

accgaagaaa ctgttcagga gtgggtggac aaaatccgta agcaggttgc cccgttcctc 300

gatctcgact gtggagaaaa ctctgctatc gcggccaata attatgactg gttcggcaat 360

atgaatgtgc tgaccttcct gcgcgatatt ggcaaacact tctccgttaa ccagatgatc 420

aacaaagaag cggttaagca gcgtctcaac cgtgaagatc aggggatttc gttcactgag 480

ttttcctaca acctgctgca gggttatagt tttgcctgtc tgaacaaaca gtacggtgtg 540

gtgctgcaaa ttggtggttc tgaccagtgg ggtaacatca cttctggtat cgacctgacc 600

cgtcgtctgc atcagaatca ggtg 624

<210> 35
<211> 624
<212> DNA
<213> artificial

<220>
<223> artificial synthetase

<400> 35
cgggggctgg tagcccaggt gacggacgag gaagcgttag cagagcgact ggcgcaaggc 60

ccgatcgcac tcacgtgtgg cttcgatcct accgctgaca gcttgcattt ggggcatctt 120

gttccattgt tatgcctgaa acgcttccag caggcgggcc acaagccggt tgcgctggta 180

ggcggcgcga cgggtctgat tggcgacccg agcttcaaag ctgccgagcg taagctgaac 240

accgaagaaa ctgttcagga gtgggtggac aaaatccgta agcaggttgc cccgttcctc 300

gatttcgact gtggagaaaa ctctgctatc gcggccaata attatgactg gttcggcaat 360

atgaatgtgc tgaccttcct gcgcgatatt ggcaaacact tctccgttaa ccagatgatc 420

aacaaagaag cggttaagca gcgtctcaac cgtgaagatc aggggatttc gttcactgag 480

ttttcctaca acctgctgca gggttatacg tttgcctgta ctaacaaaca gtacggtgtg 540

gtgctgcaaa ttggtggttc tgaccagtgg ggtaacatca cttctggtat cgacctgacc 600

cgtcgtctgc atcagaatca ggtg 624

<210> 36
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 36

Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1 5 10 15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
20 25 30

Pro Ile Ala Leu Val Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
35 40 45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
85 90 95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
100 105 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
115 120 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
130 135 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145 150 155 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
165 170 175

Leu Leu Gln Gly Tyr Ser Tyr Ala Cys Leu Asn Lys Gln Tyr Gly Val
180 185 190

```
Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
        195                 200                 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
    210                 215                 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225                 230                 235                 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
                245                 250                 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
            260                 265                 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
        275                 280                 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
    290                 295                 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305                 310                 315                 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
                325                 330                 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
            340                 345                 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
        355                 360                 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
    370                 375                 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385                 390                 395                 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
                405                 410                 415

Asn Tyr Cys Leu Ile Cys Trp Lys
            420
```

<210> 37
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 37

Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1 5 10 15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
20 25 30

Pro Ile Ala Leu Ile Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
35 40 45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
85 90 95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
100 105 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
115 120 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
130 135 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145 150 155 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
165 170 175

Leu Leu Gln Gly Tyr Ser Met Ala Cys Leu Asn Lys Gln Tyr Gly Val
180 185 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
195 200 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
210 215 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225 230 235 240

```
Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
                245                 250                 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
            260                 265                 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
        275                 280                 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
    290                 295                 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305                 310                 315                 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
                325                 330                 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
            340                 345                 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
        355                 360                 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
    370                 375                 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385                 390                 395                 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
                405                 410                 415

Asn Tyr Cys Leu Ile Cys Trp Lys
            420
```

<210> 38
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 38

```
Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1               5                   10                  15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
            20                  25                  30
```

```
Pro Ile Ala Leu Val Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
        35                  40                  45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
    50                  55                  60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65                  70                  75                  80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
                85                  90                  95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
            100                 105                 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
        115                 120                 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
    130                 135                 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145                 150                 155                 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
                165                 170                 175

Leu Leu Gln Gly Tyr Ser Met Ala Cys Ala Asn Lys Gln Tyr Gly Val
            180                 185                 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
        195                 200                 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
    210                 215                 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225                 230                 235                 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
                245                 250                 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
            260                 265                 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
        275                 280                 285
```

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
290 295 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305 310 315 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
325 330 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
340 345 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
355 360 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
370 375 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385 390 395 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
405 410 415

Asn Tyr Cys Leu Ile Cys Trp Lys
420

<210> 39
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 39

Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1 5 10 15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
20 25 30

Pro Ile Ala Leu Val Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
35 40 45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

```
Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
                85                  90                  95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
            100                 105                 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
        115                 120                 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
    130                 135                 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145                 150                 155                 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
                165                 170                 175

Leu Leu Gln Gly Tyr Ser Met Ala Cys Leu Asn Lys Gln Tyr Gly Val
            180                 185                 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
        195                 200                 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
    210                 215                 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225                 230                 235                 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
                245                 250                 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
            260                 265                 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
        275                 280                 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
    290                 295                 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305                 310                 315                 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
                325                 330                 335
```

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
340 345 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
355 360 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
370 375 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385 390 395 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
405 410 415

Asn Tyr Cys Leu Ile Cys Trp Lys
420

<210> 40
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 40

Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1 5 10 15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
20 25 30

Pro Ile Ala Leu Thr Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
35 40 45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
85 90 95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
100 105 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
115 120 125

```
Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
    130                 135                 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145                 150                 155                 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
                165                 170                 175

Leu Leu Gln Gly Tyr Thr Met Ala Cys Leu Asn Lys Gln Tyr Gly Val
            180                 185                 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
        195                 200                 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
    210                 215                 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225                 230                 235                 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
                245                 250                 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
            260                 265                 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
        275                 280                 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
    290                 295                 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305                 310                 315                 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
                325                 330                 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
            340                 345                 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
        355                 360                 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
    370                 375                 380
```

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385 390 395 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
405 410 415

Asn Tyr Cys Leu Ile Cys Trp Lys
420

<210> 41
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 41

Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1 5 10 15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
20 25 30

Pro Ile Ala Leu Thr Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
35 40 45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
85 90 95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
100 105 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
115 120 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
130 135 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145 150 155 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
165 170 175

Leu Leu Gln Gly Tyr Thr Tyr Ala Cys Leu Asn Lys Gln Tyr Gly Val
180 185 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
195 200 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
210 215 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225 230 235 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
245 250 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
260 265 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
275 280 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
290 295 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305 310 315 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
325 330 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
340 345 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
355 360 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
370 375 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385 390 395 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
405 410 415

Asn Tyr Cys Leu Ile Cys Trp Lys
420

<210> 42
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 42

Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1 5 10 15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
20 25 30

Pro Ile Ala Leu Leu Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
35 40 45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
85 90 95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
100 105 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
115 120 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
130 135 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145 150 155 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
165 170 175

Leu Leu Gln Gly Tyr Ser Met Ala Cys Ser Asn Lys Gln Tyr Gly Val
180 185 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
195 200 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
210 215 220

```
Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225                 230                 235                 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
                245                 250                 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
            260                 265                 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
        275                 280                 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
    290                 295                 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305                 310                 315                 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
                325                 330                 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
            340                 345                 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
        355                 360                 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
    370                 375                 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385                 390                 395                 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
                405                 410                 415

Asn Tyr Cys Leu Ile Cys Trp Lys
            420
```

<210> 43
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 43

```
Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1               5                   10                  15
```

```
Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
            20                  25                  30

Pro Ile Ala Leu Leu Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
        35                  40                  45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
    50                  55                  60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65                  70                  75                  80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
                85                  90                  95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
            100                 105                 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
        115                 120                 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
    130                 135                 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145                 150                 155                 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
                165                 170                 175

Leu Leu Gln Gly Tyr Ser Met Ala Cys Ala Asn Lys Gln Tyr Gly Val
            180                 185                 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
        195                 200                 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
    210                 215                 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225                 230                 235                 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
                245                 250                 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
            260                 265                 270
```

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
275 280 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
290 295 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305 310 315 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
325 330 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
340 345 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
355 360 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
370 375 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385 390 395 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
405 410 415

Asn Tyr Cys Leu Ile Cys Trp Lys
420

<210> 44
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 44

Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1 5 10 15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
20 25 30

Pro Ile Ala Leu Thr Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
35 40 45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
85 90 95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
100 105 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
115 120 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
130 135 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145 150 155 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
165 170 175

Leu Leu Gln Gly Tyr Arg Met Ala Cys Leu Asn Lys Gln Tyr Gly Val
180 185 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
195 200 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
210 215 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225 230 235 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
245 250 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
260 265 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
275 280 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
290 295 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305 310 315 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
325 330 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
340 345 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
355 360 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
370 375 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385 390 395 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
405 410 415

Asn Tyr Cys Leu Ile Cys Trp Lys
420

<210> 45
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 45

Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1 5 10 15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
20 25 30

Pro Ile Ala Leu Ile Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
35 40 45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
85 90 95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
100 105 110

```
Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
        115                 120                 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
    130                 135                 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145                 150                 155                 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
                165                 170                 175

Leu Leu Gln Gly Tyr Gly Met Ala Cys Ala Asn Lys Gln Tyr Gly Val
            180                 185                 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
        195                 200                 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
    210                 215                 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225                 230                 235                 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
                245                 250                 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
            260                 265                 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
        275                 280                 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
    290                 295                 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305                 310                 315                 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
                325                 330                 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
            340                 345                 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
        355                 360                 365
```

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
370 375 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385 390 395 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
405 410 415

Asn Tyr Cys Leu Ile Cys Trp Lys
420

<210> 46
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 46

Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1 5 10 15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
20 25 30

Pro Ile Ala Leu Gly Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
35 40 45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
85 90 95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
100 105 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
115 120 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
130 135 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145 150 155 160

```
Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
                165                 170                 175

Leu Leu Gln Gly Tyr Gly Phe Ala Cys Ala Asn Lys Gln Tyr Gly Val
            180                 185                 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
        195                 200                 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
    210                 215                 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225                 230                 235                 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
                245                 250                 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
            260                 265                 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
        275                 280                 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
    290                 295                 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305                 310                 315                 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
                325                 330                 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
            340                 345                 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
        355                 360                 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
    370                 375                 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385                 390                 395                 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
                405                 410                 415
```

Asn Tyr Cys Leu Ile Cys Trp Lys
420

<210> 47
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 47

Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1 5 10 15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
20 25 30

Pro Ile Ala Leu Gly Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
35 40 45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
85 90 95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
100 105 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
115 120 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
130 135 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145 150 155 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
165 170 175

Leu Leu Gln Gly Tyr Gly Tyr Ala Cys Met Asn Lys Gln Tyr Gly Val
180 185 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
195 200 205

```
Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
    210                 215                 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225                 230                 235                 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
                245                 250                 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
            260                 265                 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
        275                 280                 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
    290                 295                 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305                 310                 315                 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
                325                 330                 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
            340                 345                 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
        355                 360                 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
    370                 375                 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385                 390                 395                 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
                405                 410                 415

Asn Tyr Cys Leu Ile Cys Trp Lys
            420
```

<210> 48
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 48

```
Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1               5                   10                  15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
            20                  25                  30

Pro Ile Ala Leu Leu Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
        35                  40                  45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
    50                  55                  60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65                  70                  75                  80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
                85                  90                  95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
            100                 105                 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
        115                 120                 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
    130                 135                 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145                 150                 155                 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
                165                 170                 175

Leu Leu Gln Gly Tyr Ser Met Ala Cys Ala Asn Lys Gln Tyr Gly Val
            180                 185                 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
        195                 200                 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
    210                 215                 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225                 230                 235                 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
                245                 250                 255
```

```
Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
            260                 265                 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
        275                 280                 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
    290                 295                 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305                 310                 315                 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
                325                 330                 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
            340                 345                 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
        355                 360                 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
    370                 375                 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385                 390                 395                 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
                405                 410                 415

Asn Tyr Cys Leu Ile Cys Trp Lys
            420
```

<210> 49
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 49

```
Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1               5                   10                  15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
            20                  25                  30

Pro Ile Ala Leu Val Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
```

```
        35                  40                  45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
    50                  55                  60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65                  70                  75                  80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
                85                  90                  95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
            100                 105                 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
        115                 120                 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
    130                 135                 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145                 150                 155                 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
                165                 170                 175

Leu Leu Gln Gly Tyr Ser Ala Ala Cys Ala Asn Lys Gln Tyr Gly Val
            180                 185                 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
        195                 200                 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
    210                 215                 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225                 230                 235                 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
                245                 250                 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
            260                 265                 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
        275                 280                 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
    290                 295                 300
```

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305 310 315 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
325 330 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
340 345 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
355 360 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
370 375 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385 390 395 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
405 410 415

Asn Tyr Cys Leu Ile Cys Trp Lys
420

<210> 50
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 50

Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1 5 10 15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
20 25 30

Pro Ile Ala Leu Leu Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
35 40 45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr

```
                85                  90                  95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
            100                 105                 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
        115                 120                 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
    130                 135                 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145                 150                 155                 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
                165                 170                 175

Leu Leu Gln Gly Tyr Ser Ala Ala Cys Ala Asn Lys Gln Tyr Gly Val
            180                 185                 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
        195                 200                 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
    210                 215                 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225                 230                 235                 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
                245                 250                 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
            260                 265                 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
        275                 280                 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
    290                 295                 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305                 310                 315                 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
                325                 330                 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
            340                 345                 350
```

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
355 360 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
370 375 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385 390 395 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
405 410 415

Asn Tyr Cys Leu Ile Cys Trp Lys
420

<210> 51
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 51

Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1 5 10 15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
20 25 30

Pro Ile Ala Leu Val Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
35 40 45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
85 90 95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
100 105 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
115 120 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys

130 135 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145 150 155 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
165 170 175

Leu Leu Gln Gly Tyr Ser Ala Ala Cys Val Asn Lys Gln Tyr Gly Val
180 185 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
195 200 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
210 215 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225 230 235 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
245 250 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
260 265 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
275 280 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
290 295 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305 310 315 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
325 330 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
340 345 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
355 360 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
370 375 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385 390 395 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
405 410 415

Asn Tyr Cys Leu Ile Cys Trp Lys
420

<210> 52
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 52

Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1 5 10 15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
20 25 30

Pro Ile Ala Leu Ile Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
35 40 45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
85 90 95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
100 105 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asp Tyr Asp
115 120 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
130 135 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145 150 155 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
165 170 175

Leu Leu Gln Gly Tyr Asn Phe Ala Cys Val Asn Lys Gln Tyr Gly Val

```
            180                 185                 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
        195                 200                 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
    210                 215                 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225                 230                 235                 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
                245                 250                 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
            260                 265                 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
        275                 280                 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
    290                 295                 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305                 310                 315                 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
                325                 330                 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
            340                 345                 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
        355                 360                 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
    370                 375                 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385                 390                 395                 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
                405                 410                 415

Asn Tyr Cys Leu Ile Cys Trp Lys
            420
```

<210> 53
<211> 424

<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 53

Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1 5 10 15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
20 25 30

Pro Ile Ala Leu Thr Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
35 40 45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
85 90 95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
100 105 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
115 120 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
130 135 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145 150 155 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
165 170 175

Leu Leu Gln Gly Tyr Ser Ala Ala Cys Leu Asn Lys Gln Tyr Gly Val
180 185 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
195 200 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
210 215 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu

225 230 235 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
245 250 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
260 265 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
275 280 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
290 295 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305 310 315 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
325 330 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
340 345 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
355 360 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
370 375 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385 390 395 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
405 410 415

Asn Tyr Cys Leu Ile Cys Trp Lys
420

<210> 54
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 54

Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1 5 10 15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
20 25 30

Pro Ile Ala Leu Gly Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
35 40 45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
85 90 95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
100 105 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
115 120 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
130 135 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145 150 155 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
165 170 175

Leu Leu Gln Gly Tyr Ser Met Ala Cys Leu Asn Lys Gln Tyr Gly Val
180 185 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
195 200 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
210 215 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225 230 235 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
245 250 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
260 265 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu

275 280 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
290 295 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305 310 315 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
325 330 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
340 345 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
355 360 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
370 375 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385 390 395 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
405 410 415

Asn Tyr Cys Leu Ile Cys Trp Lys
420

<210> 55
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 55

Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1 5 10 15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
20 25 30

Pro Ile Ala Leu Thr Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
35 40 45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
85 90 95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
100 105 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
115 120 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
130 135 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145 150 155 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
165 170 175

Leu Leu Gln Gly Tyr Ser Ala Ala Cys Leu Asn Lys Gln Tyr Gly Val
180 185 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
195 200 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
210 215 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225 230 235 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
245 250 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
260 265 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
275 280 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
290 295 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305 310 315 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser

325 330 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
340 345 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
355 360 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
370 375 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385 390 395 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
405 410 415

Asn Tyr Cys Leu Ile Cys Trp Lys
420

<210> 56
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 56

Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1 5 10 15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
20 25 30

Pro Ile Ala Leu Ser Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
35 40 45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
85 90 95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
100 105 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
115 120 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
130 135 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145 150 155 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
165 170 175

Leu Leu Gln Gly Tyr Thr Met Ala Cys Val Asn Lys Gln Tyr Gly Val
180 185 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
195 200 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
210 215 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225 230 235 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
245 250 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
260 265 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
275 280 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
290 295 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305 310 315 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
325 330 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
340 345 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
355 360 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile

370 375 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385 390 395 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
405 410 415

Asn Tyr Cys Leu Ile Cys Trp Lys
420

<210> 57
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 57

Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1 5 10 15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
20 25 30

Pro Ile Ala Leu Ala Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
35 40 45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
85 90 95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
100 105 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
115 120 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
130 135 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145 150 155 160

```
Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
                165                 170                 175

Leu Leu Gln Gly Tyr Ser Tyr Ala Cys Leu Asn Lys Gln Tyr Gly Val
            180                 185                 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
        195                 200                 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
    210                 215                 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225                 230                 235                 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
                245                 250                 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
            260                 265                 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
        275                 280                 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
    290                 295                 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305                 310                 315                 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
                325                 330                 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
            340                 345                 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
        355                 360                 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
    370                 375                 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385                 390                 395                 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
                405                 410                 415

Asn Tyr Cys Leu Ile Cys Trp Lys
```

420

<210> 58
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 58

Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1 5 10 15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
20 25 30

Pro Ile Ala Leu Ala Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
35 40 45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
85 90 95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
100 105 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
115 120 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
130 135 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145 150 155 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
165 170 175

Leu Leu Gln Gly Tyr Thr Met Ala Cys Cys Asn Lys Gln Tyr Gly Val
180 185 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
195 200 205

```
Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
    210                 215                 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225                 230                 235                 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
                245                 250                 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
            260                 265                 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
        275                 280                 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
    290                 295                 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305                 310                 315                 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
                325                 330                 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
            340                 345                 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
        355                 360                 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
    370                 375                 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385                 390                 395                 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
                405                 410                 415

Asn Tyr Cys Leu Ile Cys Trp Lys
            420
```

<210> 59
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 59

```
Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1               5                  10                  15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
            20                  25                  30

Pro Ile Ala Leu Thr Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
        35                  40                  45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
    50                  55                  60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65                  70                  75                  80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
                85                  90                  95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
            100                 105                 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
        115                 120                 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
    130                 135                 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145                 150                 155                 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
                165                 170                 175

Leu Leu Gln Gly Tyr Thr Phe Ala Cys Met Asn Lys Gln Tyr Gly Val
            180                 185                 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
        195                 200                 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
    210                 215                 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225                 230                 235                 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
                245                 250                 255
```

```
Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
            260                 265                 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
        275                 280                 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
    290                 295                 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305                 310                 315                 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
                325                 330                 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
            340                 345                 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
        355                 360                 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
    370                 375                 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385                 390                 395                 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
                405                 410                 415

Asn Tyr Cys Leu Ile Cys Trp Lys
            420
```

<210> 60
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 60

```
Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1               5                   10                  15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
            20                  25                  30

Pro Ile Ala Leu Thr Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
        35                  40                  45
```

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
85 90 95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
100 105 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
115 120 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
130 135 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145 150 155 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
165 170 175

Leu Leu Gln Gly Tyr Ser Val Ala Cys Leu Asn Lys Gln Tyr Gly Val
180 185 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
195 200 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
210 215 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225 230 235 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
245 250 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
260 265 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
275 280 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
290 295 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305 310 315 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
325 330 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
340 345 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
355 360 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
370 375 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385 390 395 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
405 410 415

Asn Tyr Cys Leu Ile Cys Trp Lys
420

<210> 61
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 61

Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1 5 10 15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
20 25 30

Pro Ile Ala Leu Val Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
35 40 45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
85 90 95

```
Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
            100                 105                 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
        115                 120                 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
    130                 135                 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145                 150                 155                 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
                165                 170                 175

Leu Leu Gln Gly Tyr Ser Met Ala Cys Thr Asn Lys Gln Tyr Gly Val
            180                 185                 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
        195                 200                 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
    210                 215                 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225                 230                 235                 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
                245                 250                 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
            260                 265                 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
        275                 280                 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
    290                 295                 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305                 310                 315                 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
                325                 330                 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
            340                 345                 350
```

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
355 360 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
370 375 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385 390 395 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
405 410 415

Asn Tyr Cys Leu Ile Cys Trp Lys
420

<210> 62
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 62

Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1 5 10 15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
20 25 30

Pro Ile Ala Leu Ser Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
35 40 45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
85 90 95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
100 105 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
115 120 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
130 135 140

```
His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145                 150                 155                 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
                165                 170                 175

Leu Leu Gln Gly Tyr Ser Phe Ala Cys Leu Asn Lys Gln Tyr Gly Val
            180                 185                 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
        195                 200                 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
    210                 215                 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225                 230                 235                 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
                245                 250                 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
            260                 265                 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
        275                 280                 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
    290                 295                 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305                 310                 315                 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
                325                 330                 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
            340                 345                 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
        355                 360                 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
    370                 375                 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385                 390                 395                 400
```

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
405 410 415

Asn Tyr Cys Leu Ile Cys Trp Lys
420

<210> 63
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 63

Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1 5 10 15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
20 25 30

Pro Ile Ala Leu Thr Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
35 40 45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
85 90 95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
100 105 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
115 120 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
130 135 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145 150 155 160

Leu Asn Arg Glu Asp Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
165 170 175

Leu Leu Gln Gly Tyr Thr Phe Ala Cys Thr Asn Lys Gln Tyr Gly Val
180 185 190

```
Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
        195                 200                 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr
    210                 215                 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225                 230                 235                 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
                245                 250                 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
            260                 265                 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
        275                 280                 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
    290                 295                 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305                 310                 315                 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
                325                 330                 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
            340                 345                 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
        355                 360                 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
    370                 375                 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385                 390                 395                 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
                405                 410                 415

Asn Tyr Cys Leu Ile Cys Trp Lys
            420
```

<210> 64
<211> 129
<212> DNA
<213> Escherichia coli

<400> 64
agcttcccga taagggagca ggccagtaaa aagcattacc ccgtggtggg gttcccgagc 60

ggccaaaggg agcagactct aaatctgccg tcatcgacct cgaaggttcg aatccttccc 120

ccaccacca 129

<210> 65
<211> 129
<212> RNA
<213> Escherichia coli

<400> 65
agcuucccga uaagggagca ggccaguaaa aagcauuacc ccgugguggg guucccgagc 60

ggccaaaggg agcagacucu aaaucugccg ucaucgaccu cgaagguucg aauccuuccc 120

ccaccacca 129

<210> 66
<211> 34
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 66
atgaagtagc tgtcttctat cgaacaagca tgcg 34

<210> 67
<211> 34
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 67
cgaacaagca tgcgattagt gccgacttaa aaag 34

<210> 68
<211> 33
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 68
cgctactctc ccaaatagaa aaggtctccg ctg 33

<210> 69
<211> 32
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 69
ctggaacagc tatagctact gatttttcct cg 32

<210> 70
<211> 34
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 70
gccgtcacag attagttggc ttcagtggag actg 34

<210> 71
<211> 33
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 71
gattggcttc ataggagact gatatgctct aac 33

<210> 72
<211> 33
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 72
gcctctatag ttgagacagc atagaataat gcg 33

<210> 73
<211> 35
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 73
gagacagcat agatagagtg cgacatcatc atcgg 35

<210> 74
<211> 37
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 74
gaataagtgc gacatagtca tcggaagaga gtagtag 37

<210> 75
<211> 35
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 75
ggtcaaagac agttgtaggt atcgattgac tcggc 35

<210> 76
<211> 34
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 76
cgctactctc cccaaattta aaaggtctcc gctg 34

<210> 77
<211> 34
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 77
cgctactctc cccaaatata aaaggtctcc gctg 34

<210> 78
<211> 34
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 78
cgctactctc cccaaatgga aaaggtctcc gctg 34

<210> 79
<211> 34
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 79
cgctactctc cccaaagata aaaggtctcc gctg 34

<210> 80
<211> 34
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 80
cgctactctc cccaaaaaaa aaaggtctcc gctg 34

<210> 81
<211> 34
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 81
gccgtcacag atttttggc ttcagtggag actg 34

<210> 82
<211> 34
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 82
gccgtcacag attatttggc ttcagtggag actg 34

<210> 83
<211> 34
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 83
gccgtcacag attggttggc ttcagtggag actg 34

<210> 84
<211> 34
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 84
gccgtcacag atgatttggc ttcagtggag actg 34

<210> 85
<211> 34
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 85
gccgtcacag ataaattggc ttcagtggag actg 34

<210> 86
<211> 424
<212> PRT
<213> artificial

<220>
<223> artificial synthetase

<400> 86

Met Ala Ser Ser Asn Leu Ile Lys Gln Leu Gln Glu Arg Gly Leu Val
1 5 10 15

Ala Gln Val Thr Asp Glu Glu Ala Leu Ala Glu Arg Leu Ala Gln Gly
20 25 30

Pro Ile Ala Leu Ile Cys Gly Phe Asp Pro Thr Ala Asp Ser Leu His
35 40 45

Leu Gly His Leu Val Pro Leu Leu Cys Leu Lys Arg Phe Gln Gln Ala
50 55 60

Gly His Lys Pro Val Ala Leu Val Gly Gly Ala Thr Gly Leu Ile Gly
65 70 75 80

Asp Pro Ser Phe Lys Ala Ala Glu Arg Lys Leu Asn Thr Glu Glu Thr
85 90 95

Val Gln Glu Trp Val Asp Lys Ile Arg Lys Gln Val Ala Pro Phe Leu
100 105 110

Asp Phe Asp Cys Gly Glu Asn Ser Ala Ile Ala Ala Asn Asn Tyr Asp
115 120 125

Trp Phe Gly Asn Met Asn Val Leu Thr Phe Leu Arg Asp Ile Gly Lys
130 135 140

His Phe Ser Val Asn Gln Met Ile Asn Lys Glu Ala Val Lys Gln Arg
145 150 155 160

Leu Asn Arg Glu Gly Gln Gly Ile Ser Phe Thr Glu Phe Ser Tyr Asn
165 170 175

Leu Leu Gln Gly Tyr Gly Met Ala Cys Ala Asn Lys Gln Tyr Gly Val
180 185 190

Val Leu Gln Ile Gly Gly Ser Asp Gln Trp Gly Asn Ile Thr Ser Gly
195 200 205

Ile Asp Leu Thr Arg Arg Leu His Gln Asn Gln Val Phe Gly Leu Thr

210 215 220

Val Pro Leu Ile Thr Lys Ala Asp Gly Thr Lys Phe Gly Lys Thr Glu
225 230 235 240

Gly Gly Ala Val Trp Leu Asp Pro Lys Lys Thr Ser Pro Tyr Lys Phe
245 250 255

Tyr Gln Phe Trp Ile Asn Thr Ala Asp Ala Asp Val Tyr Arg Phe Leu
260 265 270

Lys Phe Phe Thr Phe Met Ser Ile Glu Glu Ile Asn Ala Leu Glu Glu
275 280 285

Glu Asp Lys Asn Ser Gly Lys Ala Pro Arg Ala Gln Tyr Val Leu Ala
290 295 300

Glu Gln Val Thr Arg Leu Val His Gly Glu Glu Gly Leu Gln Ala Ala
305 310 315 320

Lys Arg Ile Thr Glu Cys Leu Phe Ser Gly Ser Leu Ser Ala Leu Ser
325 330 335

Glu Ala Asp Phe Glu Gln Leu Ala Gln Asp Gly Val Pro Met Val Glu
340 345 350

Met Glu Lys Gly Ala Asp Leu Met Gln Ala Leu Val Asp Ser Glu Leu
355 360 365

Gln Pro Ser Arg Gly Gln Ala Arg Lys Thr Ile Ala Ser Asn Ala Ile
370 375 380

Thr Ile Asn Gly Glu Lys Gln Ser Asp Pro Glu Tyr Phe Phe Lys Glu
385 390 395 400

Glu Asp Arg Leu Phe Gly Arg Phe Thr Leu Leu Arg Arg Gly Lys Lys
405 410 415

Asn Tyr Cys Leu Ile Cys Trp Lys
420

<210> 87
<211> 6
<212> PRT
<213> artificial

<220>
<223> tryptic peptide including unnatural amino acids

<220>

<221> MISC_FEATURE
<222> (2)..(2)
<223> X is an unnatural amino acid (p-acetyl-L-phenylalanine, p-benzoyl-L-phenylalanine, p-azido-L-phenylalanine, O-methyl-L-tyrosine, or p-iodo-L-phenylalanine) or trypotophan, tyrosine, or leucine

<400> 87

Val Xaa Gly Ser Ile Lys
1 5

<210> 88
<211> 11
<212> DNA
<213> artificial

<220>
<223> B box

<220>
<221> misc_feature
<222> (8)..(8)
<223> n is a, c, g, or t

<400> 88
ggttcgantc c 11

<210> 89
<211> 82
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 89
gggggggaccg gtgggggggac cggtaagctt cccgataagg gagcaggcca gtaaaaagca 60

ttaccccgtg gtgggttccc ga 82

<210> 90
<211> 90
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 90
ggcggcgcta gcaagcttcc cgataaggga gcaggccagt aaaaagggaa gttcagggac 60

ttttgaaaaa aatggtggtg ggggaaggat 90

<210> 91
<211> 68
<212> DNA
<213> artificial

<220>

<223> oligonucleotide primer

<220>
<221> misc_feature
<222> (1)..(1)
<223> n=I

<220>
<221> misc_feature
<222> (14)..(14)
<223> n=I

<400> 91
nggggggacc ggtnggggggg accggtcggg atcgaagaaa tgatggtaaa tgaaatagga 60

aatcaagg 68

<210> 92
<211> 62
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 92
gggggggaat tcagttgatt gtatgcttgg tatagcttga aatattgtgc agaaaaagaa 60

ac 62

<210> 93
<211> 86
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 93
tcataacgag aattccggga tcgaagaaat gatggtaaat gaaataggaa atctcataac 60

gagaattcat ggcaagcagt aacttg 86

<210> 94
<211> 72
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 94
ttactacgtg cggccgcatg gcaagcagta acttgttact acgtgcggcc gcttatttcc 60

agcaaatcag ac 72

<210> 95
<211> 28
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 95
ccgatcgcgc tcgcttgcgg cttcgatc 28

<210> 96
<211> 27
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 96
atcgcggcga acgcctatga ctggttc 27

<210> 97
<211> 40
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 97
gttgcagggt tatgccgccg cctgtgcgaa caaacagtac 40

<210> 98
<211> 26
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 98
gccgctttgc tatcaagtat aaatag 26

<210> 99
<211> 21
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 99
caagccgaca accttgattg g 21

<210> 100
<211> 60
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 100

ggggacaagt ttgtacaaaa aagcaggcta cgccaatttt aatcaaagtg ggaatattgc 60

<210> 101
<211> 60
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 101
ggggacaagt ttgtacaaaa aagcaggcta ggccaatttt aatcaaagtg ggaatattgc 60

<210> 102
<211> 58
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 102
ggggaccact ttgtacaaga aagctgggtt actcttttt tgggtttggt ggggtatc 58

<210> 103
<211> 22
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 103
aagctatacc aagcatacaa tc 22

<210> 104
<211> 49
<212> DNA
<213> artificial

<220>
<223> oligonucleotide primer

<400> 104
acaaggcctt gctagcttac tctttttttg ggtttggtgg ggtatcttc 49